SEQUENCE LISTING

<110> MIROCHNITCHENKO, Oleg
WEI, Jiang
INOUYE, Masayori

<120> SOLUBLE ISCHEMIA ACTIVATED PROTEIN

<130> 266/171

<140> US 09/960,631
<141> 2001-09-20

<150> US 60/233,819
<151> 2000-09-20

<160> 8

<170> PatentIn version 3.1

<210> 1
<211> 840
<212> DNA
<213> Homo sapiens

<400> 1
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gcggcccccg gcgcccggct gttgcggctc ccggggagcg gggccgtgca ggccgcgagc 180

ccggagcgcg ccggctggac cgaggcgctg cgggccgccg tggccgagct gcgcgccggc 240

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tgcctcggcc gcgtggccga cgtctacaga tactgccgtg tgagagtacc tgaggggctc 420

ctgaaagacc tactgccagg accagtgacc ctggtgatgg aacgctcgga ggagctcaac 480

aaggacctaa acccttttac gcctcttgta ggcattcgga ttcctgatca tgcttttatg 540

caagacttgg ctcagatgtt tgagggtccg cttgctctca ctagtgccaa cctcagctcc 600

caggccagtt ctctgaatgt cgaggagttc caggatctct ggcctcagtt gtccttggtt 660

attgatgggg gacaaattgg ggatggccag agccccgagt gtcgccttgg ctcaactgtg 720

gttgatttgt ctgtgcccgg aaagtttggc atcattcgtc caggctgtgc cctggaaagt 780

actacagcca tcctccaaca gaagtacgga ctgctcccct cacatgcgtc ctacctgtga 840

<210> 2
<211> 279
<212> PRT
<213> Homo sapiens

<400> 2

Met Ser Pro Ala Arg Arg Cys Arg Gly Met Arg Ala Ala Val Ala Ala
1 5 10 15

Ser Val Gly Leu Ser Glu Gly Pro Ala Gly Ser Arg Ser Gly Arg Leu
20 25 30

Phe Arg Pro Pro Ser Pro Ala Pro Ala Ala Pro Gly Ala Arg Leu Leu
35 40 45

Arg Leu Pro Gly Ser Gly Ala Val Gln Ala Ala Ser Pro Glu Arg Ala
50 55 60

Gly Tyr Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala Gly
65 70 75 80

Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys Ala
85 90 95

Ala Ser Cys Ser Ala Ala Leu Arg Ala Val Tyr Arg Leu Lys Gly Arg
100 105 110

Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp Val
115 120 125

Tyr Arg Tyr Cys Arg Val Arg Val Pro Glu Gly Leu Leu Lys Asp Leu
130 135 140

Leu Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu Asn
145 150 155 160

Lys Asp Leu Asn Pro Phe Thr Pro Leu Val Gly Ile Arg Ile Pro Asp
165 170 175

His Ala Phe Met Gln Asp Leu Ala Gln Met Phe Glu Gly Pro Leu Ala
180 185 190

Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Asn Val Glu
195 200 205

Glu Phe Gln Asp Leu Tyr Pro Gln Leu Ser Leu Val Ile Asp Gly Gly
210 215 220

Gln Ile Gly Asp Gly Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr Val
225 230 235 240

Val Asp Leu Ser Val Pro Gly Lys Phe Gly Ile Ile Arg Pro Gly Cys
245 250 255

Ala Leu Glu Ser Thr Thr Ala Ile Leu Gln Gln Lys Tyr Gly Leu Leu
260 265 270

Pro Ser His Ala Ser Tyr Leu
275

<210> 3
<211> 1387
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(1387)
<223> The letter "n" stands for a substitution base.

<400> 3
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ggtccgcttg ctctcactag tgccaacctc agctcccagg ccagttctct gaatgtcgag 180

gagttccagg atctctggcc tcagttgtcc ttggttattg atgggggaca aattggggat 240

ggccagagcc ccgagtgtcg ccttggctca actgtggttg atttgtctgt gcccggaaag 300

tttggcatca ttcgtccagg gtgtgcctgg gaaagtacta cagccatcct ccaacagaag 360

tacggactgc tcccctcaca tgcgtcctac ctgtgaaact ctgggaagca ggaaggccca 420

agacctggtg ctggatacta tgtgtctgtc cactgacgac tgtcaaggcc tcatttgcag 480

aggccaccgg agctagggca ctagcctgac ttttaaggca gtgtgtcttt ctgagcactg 540

tagaccaagc ccttggagct gctggtttag ccttgcacct ggggaaagga tgtatttatt 600

tgtattttca tatatcagcc aaaagctgaa tggaaaagtt aagaacattc ctaggtggcc 660

ttattctaat aagtttcttc tgtctgtttt gtttttcaat tgaaaagtaa ttaaataaca 720

gatttagaat ctagtgagag cntcctctct gggggtggtg gcatttaagg ttcaacccan 780

ccnagaagtg ctgcgctgtt taaaaagtct caggtggctg cgtgtggtgg ctcatgcctg 840

taatcccaac attctgggag gcccaggcgg gagaactgct tgagcccagg agttcagaat 900

cagcctgggc aacatagcaa tactccgtct cataaaaatt aataaataaa aagtctcagg 960

tgaccaaagg ctcctgaagc tagaaccagg tttggataaa gattgaagag ccacaggcca 1020

ctcttccctc tgagccattg ggcctagtgg tgtcatgtat tgtaattgct cgcagggaga 1080

gcagtctttt tggtgtaata gtgggatgtc tgcttagttg gcaggggttc agtccaaatg 1140

gaagaatatt gggaaataaa cctccnctat cctttatagc cagggacttt tttcttattt 1200

attcataaaa taaattatag ttaattatac ccataacacc tttatttaaa tccagtgttc 1260

tccgcagcct tttgtctatt tatatgtgta ccaagtgtta aacataatta ttattgggca 1320

tttgaacntg tttttcntta naganatnct gnnattaaac atatttgtna atggnaaaaa 1380

aaaaaaa 1387

<210> 4
<211> 930
<212> DNA
<213> Mus musculus

<400> 4
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agcgacgggc cggctagttc tggccgcggc tgccgcctcc tactccctcc tgagcccgct 120

ccggcgctgc cgggggcccg gctgctgcgg cttccggaga gcgagcccgt ggaagccgcg 180

agccccgagc gcgccggctg gaccgaggcg ctgcgggccg ccgtggccga gctgcgcgcc 240

ggcgccgtgg tggcggtccc gaccgacacg ctctacggcc tggcctgctc ggcgagcagc 300

tcggcggccc tgagttgcgt gtaccgcctc aaaggccgca gcgaggccaa gccgctggcc 360

gtgtgcctgg gccgcgtggc cgacgtctac aggtactgtc aggtgagagt acctagggag 420

ctcctggaag acctgttccc aggccctgtg accctggtga tggagcgctc cgaggagctc 480

aacaaagacc tgaacccctt tactcgtctt gttggcatcc ggattcctga ccatgccttc 540

atgctggact tggcccagat gtttgggggа ccacttgcac tcactagtgc caacctcagc 600

tcccaggcca gttctctgag tgttgaggag ttccaagacc tctggcctca tttgtccctt 660

gtcattgatg gggggccaat tggggatagt cagagccctg agtgtcgcct cggctctact 720

gtggttgact tatctgtgcc tggaaagttt ggcattattc gcccaggctg tgccctggaa 780

aacactacat cgatcctcca gcagaaatat gggctgctcc cttcacaggg gtcctgttca 840

tgaaacttgg gaggacccaa ggacatgctg gatactatgt gtctgctact ggatgcaaag 900

cctcattgcc tgaggttcct acatctatag 930

<210> 5
<211> 280
<212> PRT
<213> Mus musculus

<400> 5

Met Ser Thr Ala Arg Pro Cys Ala Gly Leu Arg Ala Ala Val Ala Ala
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Gly Met Gly Leu Ser Asp Gly Pro Ala Ser Ser Gly Arg Gly Cys Arg
20 25 30

Leu Leu Leu Pro Pro Glu Pro Ala Pro Ala Leu Pro Gly Ala Arg Leu
35 40 45

Leu Arg Leu Pro Glu Ser Glu Pro Val Glu Ala Ala Ser Pro Glu Arg
50 55 60

Ala Gly Tyr Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala
65 70 75 80

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Gly Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys
                85                  90                  95

Ser Ala Ser Ser Ser Ala Ala Leu Ser Cys Val Tyr Arg Leu Lys Gly
            100                 105                 110

Arg Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp
        115                 120                 125

Val Tyr Arg Tyr Cys Gln Val Arg Val Pro Arg Glu Leu Leu Glu Asp
    130                 135                 140

Leu Phe Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu
145                 150                 155                 160

Asn Lys Asp Leu Asn Pro Phe Thr Arg Leu Val Gly Ile Arg Ile Pro
                165                 170                 175

Asp His Ala Phe Met Leu Asp Leu Ala Gln Met Phe Gly Gly Pro Leu
            180                 185                 190

Ala Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Ser Val
        195                 200                 205

Glu Glu Phe Gln Asp Leu Tyr Pro His Leu Ser Leu Val Ile Asp Gly
    210                 215                 220

Gly Pro Ile Gly Asp Ser Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr
225                 230                 235                 240

Val Val Asp Leu Ser Val Pro Gly Lys Phe Gly Ile Ile Arg Pro Gly
                245                 250                 255

Cys Ala Leu Glu Asn Thr Thr Ser Ile Leu Gln Gln Lys Tyr Gly Leu
            260                 265                 270

Leu Pro Ser Gln Gly Ser Cys Ser
        275                 280
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<210> 6
<211> 702
<212> DNA
<213> Bos taurus

<400> 6
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gggcgccgtg taccgtgtca agggccgcag cgagaccaag ccgctggccg tatgcctggg 120

ccgcgtggcc gacgtctaca ggtactgcca cgtgagagta cctgaggggc tcctgaagga 180

cctgttgcca ggaccagtga ccctggtgat ggaacgctca gaggagctca acaaggacct 240

gaatcctttc actcctcttg taggcatccg gattcctgac cacgccttca tgcaggactt 300

ggtccagatg tttgggggggc cactcgctct caccagtgcc aacctcagct cccagtccag 360

ctctctgaat gttgaggaat tccaggacct gtggcctcac ttgtccctga tcattggtgg 420

gggaccaatt ggggacggcc agagcccaga gtgtcgacta ggctcaactg tggttgactt 480

gtctgtgcct ggaaagtttg gcatcattcg tcctggttgt gcccttgaaa gtacttcagc 540

catcctccag gagtatgggc tgctcccctc acatggatcc tgctggtgac actctggagg 600

agggaaggcc caagggctgg tgctggacac tatgtgtccg actgctggtg gttggcaagg 660

cctcatttgc agaggctgct agggctacag tgttagtagt ct 702

<210> 7
<211> 126
<212> PRT
<213> Bos taurus

<400> 7

Met Glu Arg Ser Glu Glu Leu Asn Lys Asp Leu Asn Pro Phe Thr Pro
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Leu Val Gly Ile Arg Ile Pro Asp His Ala Phe Met Gln Asp Leu Val
20 25 30

Gln Met Phe Gly Gly Pro Leu Ala Leu Thr Ser Ala Asn Leu Ser Ser
35 40 45

Gln Ser Ser Ser Leu Asn Val Glu Glu Phe Gln Asp Leu Trp Pro His

50 55 60

Leu Ser Leu Ile Ile Gly Gly Gly Pro Ile Gly Asp Gly Gln Ser Pro
65 70 75 80

Glu Cys Arg Leu Gly Ser Thr Val Val Asp Leu Ser Val Pro Gly Lys
85 90 95

Phe Gly Ile Ile Arg Pro Gly Cys Ala Leu Glu Ser Thr Ser Ala Ile
100 105 110

Leu Gln Glu Tyr Gly Leu Leu Pro Ser His Gly Ser Cys Trp
115 120 125

<210> 8
<211> 841
<212> DNA
<213> Rattus novartis

<220>
<221> misc_feature
<222> (491)..(491)
<223> The letter "z" stands for sequence hybridizing.

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aaatatgggc tgctcccttc acaggggtcc tgttcatgaa acttgggagg acccaagaac 180

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ccagaagctt cgggttgagc cttgcaccca ggggaaggtt atatttactc tgtagattca 360

tgtgtcaacc cagaatggag ggaagaacat tcttagagtg accttattat tttaagtgcc 420

cctctcaccc caaccctgcc tataagttaa gtaacttgac tgcagaatta gaatgcatta 480

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gtctagaagt ctcaggacca atgcagcaaa gtctaggagc cctggccaga gctttctggg 600

tacaggagag tggtcatttg gagaaaatta ttctaggagt tccaaatgaa ataatattga 660

aaaataaaat cttgactgtt ttcagccagt gactttctta tttattggta tagttctctg 720

tttaatttat ttaactcaga agtcatcttt gttcatatgt ctacctggta tttacataat 780

tatttttaag tatttgaact gtatttcttt attaaatatt tcttctacaa aaaaaaaaaa 840

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